OTPE 463

SEQUENCE LISTING

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Gutkowska et al.
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      Disulfide bridge
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      Residues 1 and 6 are linked via a disulfide bridge
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Ala Pro Asp Leu Asp Val Arg Lys Cys Leu Pro Cys Gly Pro Gly Gly 35 40 45

Lys Gly Arg Cys Phe Gly Pro Asn Ile Cys Cys Ala Glu Glu Leu Gly 50 55 60

Cys Phe Val Gly Thr Ala Glu Ala Leu Arg Cys Gln Glu Glu Asn Tyr 65 70 75 80

Leu Pro Ser Pro Cys Gln Ser Gly Gln Lys Ala Cys Gly Ser Gly Gly 85 90 95

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The second and the second seco

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- Arg Thr Thr Arg Gln Lys His Ser Arg Leu Phe Phe Phe Met Lys His 65 70 75 80
- Leu Ser Ile Ala Asp Leu Val Val Ala Val Phe Gln Val Leu Pro Gln 85 90 95
- Leu Leu Trp Asp Ile Thr Phe Arg Phe Tyr Gly Pro Asp Leu Leu Cys
 100 105 110
- Arg Leu Val Lys Tyr Leu Gln Val Val Gly Met Phe Ala Ser Thr Tyr 115 120 125
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- Leu Arg Glu Val Ala Asp Gly Val Phe Asp Cys Trp Ala Val Phe Ile 180 185 190
- Gln Pro Trp Gly Pro Lys Ala Tyr Ile Thr Trp Ile Thr Leu Ala Val 195 200 205
- Tyr Ile Val Pro Val Ile Val Leu Ala Thr Cys Tyr Gly Leu Ile Ser 210 215 220
- Phe Lys Ile Trp Gln Asn Leu Arg Leu Lys Thr Ala Ala Ala Ala Ala 225 230 235 240
- Ala Glu Ala Pro Glu Gly Ala Ala Ala Gly Asp Gly Gly Arg Val Ala 245 250 255
- Leu Ala Arg Val Ser Ser Val Lys Leu Ile Ser Lys Ala Lys Ile Arg 260 265 270
- Thr Val Lys Met Thr Phe Ile Ile Val Leu Ala Phe Ile Val Cys Trp 275 280 285

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Pro :	Lys	Glu	Ala	Ser	Ala 310	Phe	Ile	Ile	Val	Met 315	Leu	Leu	Ala	Ser	Leu 320		
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Phe :	His	Glu	Leu 340	Val	Gln	Arg	Phe	Leu 345	Cys	Cys	Ser	Ala	Ser 350	Tyr	Leu		
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      If residue 10 not Gly, the Gly in position 9 may be optionally
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       (10)..(10)
<223>
      Xaa=Gly or nothing
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      (11)..(11)
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      If residue 10 is Gly, Xaa=Lys or nothing; if residue 10 is not Gly,
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       (12)..(12)
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Cys Tyr Ile Gln Asn Cys Pro Leu Gly Xaa Xaa Xaa